At page 72, line 7, insert --(SEQ ID NO:57)-- after "AAC-3')"
At page 72, line 8, insert --(SEQ ID NO:58)-- after "TTC-3' "
At page 73, line 7, insert --(SEQ ID NO:59)-- after "TGC-3')"
At page 73, line 14, insert --(SEQ ID NO:60)-- after "GC3' "
At page 75, line 11, insert --(SEQ ID NO:61)-- after "TAG-3' "
At page 75, line 13, insert --(SEQ ID NO:62)-- after "GGA-3' "
At page 76, line 16, insert --(SEQ ID NO:63)-- after "GAC-3')"
At page 76, line 33, insert --(SEQ ID NO:64)-- after "ACC-3')"

IN THE CLAIMS

Please amend the claims as follows:

In claim 62, delete "2" and insert -87--

In claim 63, delete "27" and insert -97--,

In claim 64, delete "28" and insert -98--

In claim 74, delete "2" and insert -87--

Add new claims 87 / 131

Delete claims 63, 64, 66-70, 72, and 73a

(Amended) An isolated DNA molecule coding for a polypeptide having the ability to

bind TNF, wherein said polypeptide is selected from the group consisting of:

A) a polypeptide comprising the amino acid sequence:

asp ser val cys pro gln gly lys tyr ile his pro gln ile his asn asn ser cys cys thr lys cys lys gly thr leu tyr tyr asn asp cys gly pro gly gln asp thr pro asp cys arg glu cys glu ser gly ser phe thr ala ser glu asn his leu his leu arg cys ser cys ser lys cys arg lys glu met gly gln val glu ile ser ser cys thr val asp thr val gly lys asp arg cys cys arg asn gln tyr his tyr glu asn leu phe gln phe arg trp ser cys asn ser leu cys asn glÿ thr val his leu ™ ser cys leu



cys	gln	glu	lys	gln	asn	thr	val	cys	thr	cys	his	ala
gly	phe	phe	leu	arg	glu	asn	glu	cys	val	ser	cys	ser
asn	cys	lys	lys	ser	leu	glu	cys	thr	lys	leu	cys	leu
pro	gln	ile	glu	asn								

[, or a C- and/or N- terminally shortened sequence thereof];

B) a polypeptide comprising the amino acid sequence:

[, or a C- and/or N- terminally shortened sequence thereof]

C) a polypeptide comprising the amino acid sequence:

asp	ser	val	cys	pro	gln	gly	lys	tyr	ile	his	pro	gln	asn
asn	ser	ile	cys	cys	thr	lys	cys	his	lys	gly	thr	tyr	leu
tyr	asn	asp	cys	pro	gly	pro	gly	gln	asp	thr	asp	cys	arg
glu	cys	glu	ser	gly	ser	phe	thr	ala	ser	glu	asn	his	leu
arg	his	cys	leu	ser	cys	ser	lys	cys	arg	lys	glu	met	gly
gln	val	glu	ile	ser	ser	cys	thr	val	asp	arg	asp	thr	val
cys	gly	cys	arg	lys	asn	gln	tyr	arg	his	tyr	trp	ser	glu
asn	leu	phe	gln	cys	phe	asn	cys	ser	leu	cys	leu	asn	gly
thr	val	his	leu	ser	cys	gln	glu	lys	gln	asn	thr	val	cys
thr	cys	his	ala	gly	phe	phe	leu	arg	glu	asn	glu	cys	val
ser	cys	ser	asn	cys	lys	lys	ser	leu	glu	cys	thr	lys	leu
cys	leu	pro	gln	ile	glu	asn	val	lys	gly	thr	glu	asp	ser
gly	thr	thr										_	

[, or a C- and/or N- terminally shortened sequence thereof]; and-

D) a polypeptide comprising the amino acid sequence:

leu val pro his leu gly asp arg glu lys arg asp ser val

1

trapp.

his pro gln gly lys tyr ile pro gln asn asn ile cys ser his cys cys thr lys cys lys gly thr tyr leu tyr asn asp thr cys pro gly pro gly gln asp asp cys arg glu cys glu ser phe thr ala ser glu his leu his gly ser asn arg cys glu leu ser ser lys cys arg lys glu met gly gln val cys ile ser ser cys thr val asp arg asp thr val cys gly cys lys gln his glu leu phe asn tyr arg tyr trp ser asn arg his gln phe asncys ser leu leu thr val cys cys asn gly leu ser cys gln glu lys gln asn thr val cys thr cys his ala gly phe phe leu arg glu glu val ser ser asn cys cys leu glu leu asn cys lys lys ser cys thr lys leu cys pro thr; gln ile glu asn val lys gly thr glu asp ser gly thr

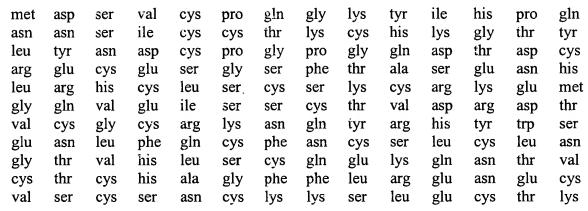
[. or a C- and/or N- terminally shortened sequence thereof]

- E) a polypeptide comprising the amino acid sequence of A, B, C or D with at least one conservative amino acid substitution;
- F) a polypeptide comprising the amino acid sequence of A, B, C or D with at least one amino acid substitution at a glycosylation site;
- G) a polypeptide comprising the amino acid sequence of A, B, C or D with at least one amino acid substitution at a proteolytic cleavage site; and
- H) a polypeptide comprising the amino acid sequence of A, B, C or D with at least one amino acid substitution at a cysteine residue.

(Amended) An isolated DNA molecule coding for a polypeptide having the ability to

bind TNF selected from the group consisting of:

A) a polypeptide comprising the amino acid sequence:



leu cys leu pro gln ile glu asn

[, or a C- and/or N- terminally shortened sequence thereof];

B) a polypeptide comprising the amino acid sequence:

met	leu	vai	pro	his	leu	gly	asp	arg	glu	lys	arg	asp	ser
val	cys	pro	gln	gly	lys	tyr	ile	his	pro	gln	asn	asn	ser
ile	cys	cys	thr	lys	cys	his	lys	gly	thr	tyr	leu	tyr	asn
asp	cys	pro	gly	pro	gly	gln	asp	thr	asp	cys	arg	glu	
-	-			•		_			-	-		_	cys
glu	ser	gly	ser	phe	thr	ala	ser	glu	asn	his	leu	arg	his
cys	leu	ser	cys	ser	lys	cys	arg	iys	glu	met	gly	gln	val
glu	ile	ser	ser	cys	thr	val	asp	arg	asp	thr	val	cys	gly
cys	arg	lys	asn	gln	tyr	arg	his	tyr	trp	ser	glu	asn	leu
phe	gln	cys	phe	asn	cys	ser	leu	cys	leu	asn	gly	thr	val
his	leu	ser	cys	gln	glu	lys	gĺn	asn	thr	val	cys	thr	cys
his	ala	gly	phe	phe	leu	arg	glu	asn	glu	cys	val	ser	cys
ser	asn	cys	lys	lys	ser	leu	glu	cys	thr	lys	leu	cys	leu
pro	gln	ile	glu	asn									

[, or a C- and/or N- terminally shortened sequence thereof];

C) a polypeptide comprising the amino acid sequence:

[, or a C- and/or N- terminally shortened sequence thereof];

D) a polypeptide comprising the amino acid sequence:

val	cys	pro	gln	gly	lys	tyr	ile	his	pro	gln	arg asn leu	asn	ser
						7							

(alle

Noort

gly asp cys pro gly pro gln asp thr asp cys arg glu cys glu thr ser gly ser phe ala ser glu asn his leu his arg cys leu ser ser lys cys lys cys arg glu met gly gln val glu ile ser ser cys thr val asp arg asp thr val cys gly cys arg lys gln asn tyr his glu arg tyr trp ser asn leu gln phe cys phe asn cys ser leu cys leu asn gly thr val his leu ser cys gln glu lys gln asn thr val thr cys cys his ala gly phe phe leu arg glu asn glu val cys ser cys ser asn cys lys lys ser leu glu thr cys lys leu cys leu pro gln ile glu asn val lys gly thr glu asp ser gly thr thr

[, or a C- and/or N- terminally shortened sequence thereof];

E) a polypeptide comprising the amino acid sequence:

thr met gly leu ser val pro asp leu leu leu pro leu val leu leu glu leu leu val gly ile tyr ser gly val ile pro gly leu his leu val pro gly asp arg glu lys arg asp ser val cys gln gly lys ile his pro tyr pro gln asn ser asn ile cys cys thr lys cys his lys gly thr tyr leu tyr asn asp cys pro gly pro gly gln asp thr asp cys arg glu cys glu ser gly ser phe thr ala ser glu his leu his asn arg leu cys ser cys ser lys cys lys glu gly gln val arg met glu ile ser ser cys thr val asp arg asp thr val gly cys cys lys arg asn gln tyr arg his glu leu tyr trp ser asn phe gln cys phe asn cys ser leu cys leu asn gly thr val his leu ser cys gln glu lys gin asn thr val cys thr cys his ala gly phe phe leu arg glu asn glu cys val ser cys ser asncys lys lys ser leu glu cys thr lys leu cys leu ile gln pro glu asn

[, or a C- and/or N- terminally shortened sequence thereof];

F) a polypeptide comprising the amino acid sequence:

met gly thr pro leu ser val pro leu leu leu leu val asp leu leu glu leu leu val gly ile tyr pro ser gly val ile gly leu val his leu glu pro gly asp arg lys arg asp ser val cys gln gly ile his pro lys tyr pro gln asn asn ser ile thr lys his cys cys cys lys gly thr tyr leu tyr asn asp gln thr cys pro gly pro gly asp asp cys arg glu cys glu ser gly ser phe thr ala his leu ser glu asn arg his cys leu ser cys ser lys cys lys gly arg glu met gln val

all of

glu cys phe his his ser	ile arg gln leu ala asn	ser lys cys ser gly cys	ser asn phe cys phe lys	cys gln asn gln phe lys	thr tyr cys glu leu ser	val arg ser lys arg leu	his leu gln glu glu	asn cys	glu thr	thr ser asn val cys lys	val glu gly cys val leu	cys asn thr thr ser cys	gly leu val cys cys leu
pro thr	gln	ile	glu	asn	val		gly	thr	glu	asp	ser	gly	thr

[, or a C- and/or N- terminally shortened sequence thereof];

G) a polypeptide comprising the amino acid sequence:

met	gly	leu	ser	thr	val	pro	asp	leu	leu	leu	pro
leu	val	leu	leu	glu	leu	leu	val	gly	ile	tyr	pro
ser	gly	val	ile	gly	asp	ser	val	-cys	pro	gln	gly
lys	tyr	ile	his	pro	gln	asn	asn	ser	ile	cys	cys
thr	lys	cys	his	lys	gly	thr	tyr	leu	tyr	asn	asp
cys	pro	gly	pro	gly	gln	asp	thr	asp	cys	arg	glu
cys	glu	ser	gly	ser	phe	thr	ala	ser	glu	asn	his
leu	arg	his	cys	leu	ser	cys	ser	iys	cys	arg	lys
glu	met	gly	gln	val	glu	ile	ser	ser	cys	thr	val
asp	arg	asp	thr	val	cys	gly	cys	arg	lys	asn	gln
tyr	arg	his	tyr	trp	ser	glu	asn	leu	phe	gln	cys
phe	asn	cys	ser	leu	cys	leu	asn	gly	thr	val	his
leu	ser	cys	gln	glu	lys	gln	asn	thr	val	cys	thr
cys	his	ala	gly	phe	phe	leu	arg	glu	asn	glu	cys
val	ser	cys	ser	asn	cys	lys	lys	ser	leu	glu	cys
thr	lys	leu	cys	leu	pro	gln	ile	glu	asn		

[, or a C- and/or N- terminally shortened sequence thereof];

H) a polypeptide comprising the amino acid sequence:

phe cys thr cys his ala gly phe leu arg glu glu asn cys val lys lys ser cys ser asn cys leu glu thr ser cys lys leu cys leu pro gln ile glu asn val lys gly thr glu asp ser gly thr thr

[, or a C- and/or N- terminally shortened sequence thereof]; and

I) a polypeptide comprising the amino acid sequence:

T

thr met gly leu ser val pro ieu leu leu leu val asp pro leu leu leu glu leu val gly ile ile tyr pro ser gly val gly leu val his leu pro gly asp arg glu lys arg asp ser val gly his cys pro gln lys tyr ile pro gln asn asn ser ile thr lys cys his lys thr leu cys cys gly tyr tyr asn asp cys gly pro gly gln asp thr cys glu cys pro. asp arg glu phe thr ala glu ser gly ser ser asn his leu arg his cys leu ser lys lys cys ser cys arg glu met gly gln val glu ile cys thr val thr val ser ser asp arg asp cys gly cys arg lys asn gln tyr arg his trp ser glu asn leu tyr phe gln phe leu val cys asn cys ser cys leu asn gly thr his leu ser cys gln glu lys gln asn thr val cys thr cys his ala gly phe phe leu glu val arg asn glu cys ser cys lys ser leu glu thr leu ser asn cys lys cys lys leu cys pro gln ile glu val lys gly thr glu thr asn asp ser gly thr val leu leu val ile leu pro phe phe gly leu cys leu leu leu leu phe ile gly leu met ser tyr gln arg arg tyr trp lys ser lys leu tyr ser ile val cys gly lys ser thr pro glu lys glu glu leu glu thr thr gly gly thr lys pro leu ala pro asn pro ser phe ser thr pro gly phe thr pro pro thr leu gly phe ser pro val pro ser ser thr phe thr ser ser ser thr tyr thr pro gly phe ala asp cys pro asn glu ala val ala gln gly ala pro arg arg pro pro tyr asp ile ala thr ala pro pro leu leu ala ser asp ile pro asn pro leu gln lys trp glu asp ser ala his lys pro gln ser leu thr ala thr leu ala glu asp asp asp pro tyr val val leu asn val pro pro leu arg trp lys glu phe val arg arg gly leu his glu ile leu gln ser asp leu glu asn asp arg gly leu glu ala gln met leu ala thr arg cys arg tyr ser thr glu ala thr leu leu trp arg arg arg pro arg arg glu leu gly val leu arg asp met asp leu leu gly cys leu arg glu asp ile glu glu ala leu cys gly pro ala ala leu pro pro ala leu leu pro ser arg;

[, or a C- and/or N- terminally shortened sequence thereof]

- a polypeptide comprising the amino acid sequence of A, B, C, D, E, F, G, H or I with at least one conservative amino acid substitution;
- a polypeptide comprising the amino acid sequence of A, B, C, D, E, F, G, H or I with at least one amino acid substitution at a glycosylation site;
- with at least one amino acid substitution at a proteolytic cleavage site; and
- 工 (作为) a polypeptide comprising the amino acid sequence of A, B, C, D, E, F, G, H or I with at least one amino acid substitution at a cysteine residue.

(Amended) An isolated DNA molecule [according to claim 87], wherein said

DNA is selected from the group consisting of:

1/1

A) a DNA comprising the sequence:

CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGC AGG AAG AAC CAC TGC CTC AGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC AAT TGC AGC CTC TGC CAG AAC CTC TGC CAG AAA CCC GTG TGC TGC ACA GTG CAC CTC TGC CAG AAA TGC CAG AAA CAG AAA CAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA AAA CAG AAA AAC CTG TGC TCC TGC CAG AAG AAC CAG AAA AAC CTG TGC TCC TGC CAG AAA TTC TTC TTT CTA AGA AAA AGC CTG GAG TGC TCC TGC CAG AAG AAA AAC CTG TGC TTC AAT TGC AAA AAC CTG TGC CAG AAA AAC CTG TGC CTA CCC CAG ATT GAG AAA

[, or a C- and/or N- terminally shortened sequence thereof];

B) a DNA comprising the sequence:

CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG

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GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC AAT TGC AGC CTC TGC CTC AAT GGG ACC GTG CAC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AGT AAC TGT AAG AAA AGC CTG GAG TGC ACG AAG TTG TGC CTA CCC CAG ATT GAG AAT GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA
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[, or a C- and/or N- terminally shortened sequence thereof];

C) A DNA comprising the sequence:
GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA
AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC
TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG
GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA
GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC
CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA
GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG
TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC
AAT TGC AGC CTC TGC CTC AAT GGG ACC GTG CAC CTC TCC
TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA
GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AGT
AAC TGT AAG AAA AGC CTG GAG TGC ACG AAG TTG TGC CTA

[, or a C- and/or N- terminally shortened sequence thereof]; and a DUA molecule

D) -DNA comprising the sequence:

CCC CAG ATT GAG AAT

GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC AAT TGC AGC CTC AAT GGG ACC GTG TGC CAC CTC TCC TGC CAG GAG AAA CAC GTG TGC ACC GTG TGC ACC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AGT AAC TGT AAG AAA AGC CTG GAG TGC ACC TGC CAT GCA CCC CAG ATT GAG AAT GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA;

[, or a C- and/or N- terminally shortened sequence thereof]

(950x

molecule comprising the a DNA sequence of A, B, C or D encoding at least one conservative amino acid substitution: molecule comprising the a DNA sequence of A, B, C or D encoding at least one amino acid substitution at a glycosylation site: moderate comprising the a DNA sequence of A, B, C or D encoding at least one amino acid substitution at a proteolytic cleavage site; and a DNA sequence of A, B, C or D encoding at least one amino acid substitution at a cysteine residue. An isolated DNA molecule (Amended) DNA coding for a polypeptide having the ability to bind to TNF, wherein said DNA coding said polypeptide is selected from the group consisting of: a DDA molecule DNA comprising the sequence: ATG CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC AAT TGC AGC CTC TGC CTC AAT GGG ACC GTG CAC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AGT AAC TGT AAG AAA AGC CTG GAG TGC ACG AAG TTG TGC CTA CCC CAG ATT GAG AAT [, or a C- and/or N- terminally shortened sequence thereof]; a DUA molecule **DNA** comprising the sequence: ATG CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA 13

AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC AAT TGC AGC CTC TGC CAC GTG CAC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AGT AAC TGT AAG AAA AGC CTG GAG TGC ACC ACG ATT GAG AAT GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA

[, or a C- and/or N- terminally shortened sequence thereof];

C) -DNA comprising the sequence:

[, or a C- and/or N- terminally shortened sequence thereof];

D) DNA comprising the sequence:

ATG GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAC TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TCC AAT TGC AAT TGC AGC GTG TGC ACC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC CTC TCC TGC CAG GAG AAA CAG AAC GAG TGT TTC TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGC CAT

AGT AAC TGT AAG AAA AGC CTG GAG TGC ACG AAG TTG TGC CTA CCC CAG ATT GAG AAT GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA

[, or a C- and/or N- terminally shortened sequence thereof];

a DNA molecule

E) DNA comprising the sequence:

ATG GGC CTC TCC ACC GTG CCT GAC CTG CTG CCA CTG GTG CTC CTG GAG CTG TTG GTG GGA ATA TAC CCC TCA GGG GTT ATT GGA CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC AAT TGC AGC CTC TGC CTC AAT GGG ACC GTG CAC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AGT AAC TGT AAG AAA AGC CTG GAG TGC ACG AAG TTG TGC CTA CCC CAG ATT GAG AAT

[, or a C- and/or N- terminally shortened sequence thereof];

F) A DNA comprising the sequence:

CTA CCC CAG ATT GAG AAT GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA

[, or a C- and/or N- terminally shortened sequence thereof];

G) / DNA comprising the sequence:

ATG GGC CTC TCC ACC GTG CCT GAC CTG CTG CCA CTG GTG CTC CTG GAG CTG TTG GTG GGA ATA TAC CCC TCA GGG GTT ATT GGA GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC AAT TGC AGC CTC TGC CTC AAT GGG ACC GTG CAC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AGT AAC TGT AAG AAA AGC CTG GAG TGC ACG AAG TTG TGC CTA CCC CAG ATT GAG AAT

[, or a C- and/or N- terminally shortened sequence thereof];

H) A DNA comprising the sequence:

ATG GGC CTC TCC ACC GTG CCT GAC CTG CTG CCA CTG GTG CTC CTG GAG CTG TTG GTG GGA ATA TAC CCC TCA GGG GTT ATT GGA GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC AAT TGC AGC CTC TGC CTC AAT GGG ACC GTG CAC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AGT AAC TGT AAG AAA AGC CTG GAG TGC ACG AAG TTG TGC CTA CCC CAG ATT GAG AAT GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA

T I

(agly

[, or a C- and/or N- terminally shortened sequence thereof]; and a DNA molecule

I) A DNA comprising the sequence:

ATG GGC CTC TCC ACC GTG CCT GAC CTG CTG CCA CTG GTG CTC CTG GAG CTG TTG GTG GGA ATA TAC CCC TCA GGG GTT ATT GGA CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC AAT TGC AGC CTC TGC CTC AAT GGG ACC GTG CAC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AGT AAC TGT AAG AAA AGC CTG GAG TGC ACG AAG TTG TGC CTA CCC CAG ATT GAG AAT GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA GTG CTG TTG CCC CTG GTC ATT TTC TTT GGT CTT TGC CTT TTA TCC CTC CTC TTC ATT GGT TTA ATG TAT CGC TAC CAA CGG TGG AAG TCC AAG CTC TAC TCC ATT GTT TGT GGG AAA TCG ACA CCT GAA AAA GAG GGG GAG CTT GAA GGA ACT ACT ACT AAG CCC CTG GCC CCA AAC CCA AGC TTC AGT CCC ACT CCA GGC TTC ACC CCC ACC CTG GGC TTC AGT CCC GTG CCC AGT TCC ACC TTC ACC TCC AGC TCC ACC TAT ACC CCC GGT GAC TGT CCC AAC TTT GCG GCT CCC CGC AGA GAG GTG GCA CCA CCC TAT CAG GGG GCT GAC CCC ATC CTT GCG ACA GCC CTC GCC TCC GAC CCC ATC CCC AAC CCC CTT CAG AAG TGG GAG GAC AGC GCC CAC AAG CCA CAG AGC CTA GAC ACT GAT GAC CCC GCG ACG CTG TAC GCC GTG GTG GAG AAC GTG CCC CCG TTG CGC TGG AAG GAA TTC GTG CGG CGC CTA GGG CTG AGC GAC CAC GAG ATC GAT CGG CTG GAG CTG CAG AAC GGG CGC TGC CTG CGC GAG GCG CAA TAC AGC ATG CTG GCG ACC TGG AGG CGG CGC ACG CCG CGC GAG GCC ACG CTG GAG CTG CTG GGA CGC GTG CTC CGC GAC ATG GAC CTG CTG GGC TGC CTG GAG GAC ATC GAG GAG GCG CTT TGC GGC CCC GCC GCC CTC CCG CCC GCG CCC AGT CTT CTC AGA,

[, or a C- and/or N- terminally shortened sequence thereof]

J) a DNA sequence of A, B, C, D, E, F, G, H or I encoding at least one conservative amino acid substitution;

indecide compersing the

- a DNA sequence of A, B, C, D, E, F, G, H or I encoding at least one amino acid substitution at a glycosylation site;
 - moderale comparsing the a DNA, sequence of A, B, C, D, E, F, G, H or I encoding at least one amino acid substitution at a proteolytic cleavage site; and

motecule comprising the

- a DNA sequence of A, B, C, D, E, F, G, H or I encoding at least one amino acid substitution at a cysteine residue.
- recombinant (Amended) A recombinant host cell containing a DNA molecule comprising a V DNA coding for a polypeptide having the ability to bind TNF selected from the group consisting of:
 - A) a polypeptide comprising the amino acid sequence:

met gly leu ser thr val pro asp leu leu leu leu val pro leu leu leu leu val ile glu gly tyr pro ser gly val ile gly leu val his leu glu pro gly asp arg lys ser arg asp val cys pro gln gly lys tyr ile his pro gln asn asn ser ile thr lys his gly thr cys cys cys lys tyr leu tyr asn gly gln thr asp cys pro gly pro asp asp cys arg glu cys glu phe thr ala glu his leu his ser gly ser ser asn arg cys leu ser cys ser lys cys arg lys glu met gly gln val glu ile ser thu val thr val \cdot gly ser cys asp asp cys arg cys arg lys asn gln tyr arg his tyr trp ser glu asn leu phe gln cys phe asn cys ser leu leu thr val cys asn gly his leu gln ser cys gln glu lys asn thr val cys thr cys his ala gly phe phe leu arg glu asn glu cys val ser cys ser asn cys lys lys ser leu glu cys thr lys leu leu cys gln ile glu val lys gly pro asn thr glu asp gly thr ser thr val. leu leu leu val ile leu pro phe phe gly leu cys leu ser leu leu phe ile gly leu met tyr arg tyr gln arg trp lys ser lys leu tyr ser ile val gly lys ser thr cys lys leu pro glu glu gly glu glu gly thr thr thr lys pro leu ala phe pro asn pro ser ser thr gly phe thr pro pro thr leu phe ser val thr phe pro gly pro pro ser ser thr ser ser ser thr tyr thr pro gly phe ala asp cys pro asn ala pro arg glu val ala pro gln gly ala arg pro tyr asp pro ile leu ala thr ala leu ala ser asp pro ile pro asn pro leu gln lys trp glu ser ala his lys gln asp pro ser leu asp thr pro ala thr leu ala val val glu asp asp tyr asn val leu trp lys glu phe val leu pro pro arg arg arg gly leu ser asp his glu ile asp arg leu glu leu gln asn

- PORTOR

gly	arg	cys	leu	arg	glu	ala	gln	tyr	ser	met	leu	ala	thr
trp	arg	arg	arg	thr	pro	arg	arg	glu	ala	thr	leu	glu	leu
leu	gly	arg	val	leu	arg	asp	met	asp	leu	leu	gly	cys	leu
glu	asp	ile	glu	glu	ala	leu	cys	gly	pro	ala	ala	leu	pro
pro	ala	pro	ser	leu	leu	arg;	-		_				•

B) a polypeptide comprising the amino acid sequence:

asp	ser	val	cys	pro	gln	gly	lys	tyr	ile	his	pro	gln	asn
asn	ser	ile	cys	cys	thr	lys	cys	his	lys	gly	thr	tyr	leu
tyr	asn	asp	cys	pro	gly	pro	gly	gln	asp	thr	asp	cys	arg
glu	cys	glu	ser	gly	ser	phe	thr	ala	ser	glu	asn	his	leu
arg	his	cys	leu	ser	cys	ser	lys	cys	arg	lys	glu	met	gly
gln	val	glu	ile	ser	ser	cys	thr	val	asp	arg	asp	thr	val
cys	gly	cys	arg	lys	asn	gln	tyr	arg	his	tyr	trp	ser	glu
asn	leu	phe	gln	cys	phe	asn	cys	ser	leu	cys	leu	asn	gly
thr	val	his	leu	ser	cys	gln.	glu	lys	gln	asn	thr	val	cys
thr	cys	his	ala	gly	phe	phe	leu	arg	glu	asn	glu	cys	val
ser	cys	ser	asn	cys	lys	lys	ser	leu	glu	cys	thr	lys	leu
cys	leu	pro	gln	ile	glu	asn •	/ [: and	d					

- C) a fragment of A or B]
- C) a polypeptide comprising the amino acid sequence of A, B, C or D with at least one conservative amino acid substitution;
- D) a polypeptide comprising the amino acid sequence of A, B, C or D with at least one amino acid substitution at a glycosylation site;
- E) a polypeptide comprising the amino acid sequence of A, B, C or D with at least one amino acid substitution at a proteolytic cleavage site; and
- F) a polypeptide comprising the amino acid sequence of A, B, C or D with at least one amino acid substitution at a cysteine residue.

(Amended) A process according to claim 78, wherein the DNA molecule comprises promoter DNA, other than the promoter DNA for the native polypeptide having TNF inhibitory activity, operatively linked to the nucleic acid encoding the [TNF inhibitor] native polypeptide having TNF inhibitory activity.

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An isolated DNA molecule wherein said DNA comprises a sequence selected from

the group consisting of:

A)

R² GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CQA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TCT TGC ACA GTG GAC CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TCC AAT TGC CAG GAG AAA CCTC TGC CTC AAT GGG ACC GTG CAC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AAT AAC TGT AAG AAA AGC CTG GAG TGC ACC AAG TTC TTC TGT AAT CCC CAG ATT GAG AAA

wherein R² is absent or is a DNA comprising a sequence coding for a polypeptide which can be 7cleaved in vivo;

- B) a fragment or degenerate variant of the polypeptide of A;
- C) a polypeptide comprising the amino acid sequence of A or B with at least one conservative amino acid substitution;
- D) a polypeptide comprising the amino acid sequence of A or B with at least one amino acid substitution at a glycosylation site;
- E) a polypeptide comprising the amino acid sequence of A or B with at least one amino acid substitution at a proteolytic cleavage site; and
- F) a polypeptide comprising the amino acid sequence of A or B with at least one amino acid substitution at a cysteine residue.

An isolated DNA molecule according to claim of, wherein R² is a DNA comprising a sequence which codes entirely or partly for a signal sequence.

An isolated DNA molecule according to claim -87, wherein R² is a DNA molecule comprising the sequence CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA or a fragment thereof.

An isolated DNA molecule according to claim 38, wherein R² is a DNA molecule comprising the sequence R³ CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA, wherein R³ is a DNA coding for a signal peptide, or a fragment thereof.

An isolated DNA molecule according to claim 98, wherein R³ is a DNA molecule comprising the sequence:

ATG GGC CTC TCC ACC GTG CCT GAC CTG CTG CTG CCA CTG GTG CTC CTG GAG CTG TTG GTG GGA ATA TAC CCC TCA GGG GTT ATT GGA, or a fragment thereof.

A nucleic acid which hybridizes with DNA complementary to the DNA defined in claim 27 under conditions of low stringency such that the nucleic acid is useful as a hybridization probe to detect DNA encoding the polypeptide of A or B.

A recombinant DNA molecule, which is replicable in prokaryotic or eukaryotic host organisms, wherein said DNA molecule contains expression control sequences functionally linked to the DNA sequence defined in claim 87, or a degenerate variant or a fragment thereof.

A process for preparing a recombinant TNF receptor protein, comprising cultivating the host cell of claim 114 and isolating the expressed protein.

An isolated DNA molecule coding for a polypeptide selected from the group consisting of:

val

pro

asp

leu

gly

cys

lys

leu

leu

tyr

gly

gly

cys

asp

glu

his

lys

val

gln

cys

his

thr

cys

cys

lys

pro

leu

trp

ser

pro

pro

asp

lys

thr

cys

cys

leu

glu

asp

tyr

phe

leu

cys

val

thr:

gly :

leu

leu

lys

thr

A) a polypeptide comprising the amino acid sequence:

ser

thr

met

ser

lys

leu

tyr

gly

leu

110204 leu val leu leu glu leu leu val gly ile ser gly val ile gly leu his val leu pro glu arg lys arg asp ser val cys pro gln tyr ile his pro gln asn asn ser ile cys lys his cys lys gly thr tyr leu tyr asn pro gly pro gly gln asp thr asp cys arg glu ser gly ser thr phe ala ser glu asn arg his leu cys ser cys ser lys cys arg met gly gln val glu ile ser ser cys thr arg asp thr val cys gly lys .cys arg asn arg his trp tyr ser glu asn leu phe gln leu asn cys ser leu thr cys asn gly val ser cys glu gln lys gln thr val asn cys his ala gly phe phe leti glu arg asn glu ser cys ser asn cys lys lys ser leu glu lys leu cys leu gln ile pro glu asn val thr glu asp ser gly thr thr val leu leu ile val phe phe gly leu cys leu leu ser phe ile leu gly met tyr gln arg tyr arg

ser

ile

pro glu lys glu glu leu gly gly glu thr thr thr lys leu ala pro pro asn pro ser phe thr ser pro pro gly phe thr pro thr leu gly phe val ser pro ser ser thr phe thr pro ser ser ser thr tyr thr gly pro cys asn ala ala asp pro phe pro arg arg glu val ala pro pro tyr gln gly ala asp pro ile leu ala thr ala leu ala ile ser asp pro pro asn pro leu gln lys glu ala his lys trp asp ser pro gln ser leu thr asp asp asp ala thr leu tyr pro ala val val glu asn val leu pro pro arg trp lys glu phe val leu arg arg gly leu ser his asp glu ile asp arg leu glu leu gln gly leu asn arg cys

val

arg	glu	ala	gln	tyr	ser	met	leu	ala	thr	trp	arg
arg	arg	thr	pro	arg	arg	glu	ala	thr	leu	glu	leu
leu	gly	arg	val	leu	arg	asp	met	asp	leu	leu	gly
cys	leu	glu	asp	ile	glu	glu	ala	leu	cys	gly	pro
ala	ala	leu	pro	pro	ala	pro	ser	leu	leu	arg;	_

B) a polypeptide comprising the amino acid sequence:

asp	ser	val	cys	pro	gln	gly	lys	tyr	ile	his	pro
gln	asn	asn	ser	ile	cys	cys	thr	lys	cys	his	lys
gly	thr	tyr	leu	tyr	asn	asp	cys	pre	gly	pro	gly
gln	asp	thr	asp	cys	arg	glu	cys	glu	ser	gly	ser
phe	thr	ala	ser	glu	asn	his	leu	arg	his	cys	leu
ser	cys	ser	lys	cys	arg	lys	glu	met	gly	gln	val
glu	ile	ser	ser	cys	thr	val	asp	arg	asp	thr	val
cys	gly	cys	arg	lys	asn	gln	tyr	arg	his	tyr	trp
ser	glu	asn	leu	phe	gln	cys	phe	asn	cys	ser	leu
cys	leu	asn	gly	thr	val	his	leu	ser	cys	gln	glu
lys	gln	asn	thr	val	cys	thr	cys	his	ala	gly	phe
phe	leu	arg	glu	asn	glu	cys	val	ser	cys	ser	asn
cys	lys	lys	ser	leu	glu	cys	thr	lys	leu	cys	leu
pro	gln	ile	glu	asn;	[and]						

- C) a fragment of A or B complementary to the DNA encoding the polypeptide of A or B and is useful as a hybridization probe to detect the DNA encoding the polypeptide of A or B;
- D) a polypeptide comprising the amino acid sequence of A, B or C with at least one conservative amino acid substitution;
- E) a polypeptide comprising the amino acid sequence of A, B or C with at least one amino acid substitution at a glycosylation site;
- F) a polypeptide comprising the amino acid sequence of A, B or C with at least one amino acid substitution at a proteolytic cleavage site; and
- G) a polypeptide comprising the amino acid sequence of A, B or C with at least one amino acid substitution at a cysteine residue.

A DNA according to claim 95, wherein said polypeptide is selected from the group consisting of:

A) a polypeptide comprising the amino acid sequence:

\mathbb{R}^2	asp	ser	val	cys	pro	gln	gly	lys	tyr	ile	his
pro	gln	asn	asn	ser	ile	cys	cys	thr	lys	cys	his
lys	gly	thr	tyr	leu	tyr	asn	asp	cys	pro	gly	pro
gly	gln	asp	thr	asp	cys	arg	glu	cys	glu	ser	gly
ser	phe	thr	ala	ser	glu	asn	his	leu	arg	his	cys
leu	ser	cys	ser	lys	cys	arg	lys	glu	met	gly	gln
val	glu	ile	ser	ser	cys	thr	val	asp	arg	asp	thr
val	cys	gly	cys	arg	lys	asn	gln	tyr	arg	his	tyr
trp	ser	glu	asn	leu	phe	gln	cys	phe	asn	cys	ser
leu	cys	leu	asn	gly	thr	val	his	leu	ser	cys	gln
glu	lys	gln	asn	thr	val	cys	thr	cys	his	ala	gly
phe	phe	leu	arg	glu	asn	glu	cys	val	ser	cys	ser
asn	cys	lys	lys	ser	leu	glu	cys	thr	lys	leu	cys
leu	pro	gln	ile	glu	asnį						

wherein R² is absent or is a polypeptide which can be cleaved in vivo;

- B) a fragment or functional derivative of the polypeptide of A which binds TNF;
- C) a polypeptide comprising the amino acid sequence of A or B with at least one conservative amino acid substitution;
- D) a polypeptide comprising the amino acid sequence of A or B with at least one amino acid substitution at a glycosylation site;
- E) a polypeptide comprising the amino acid sequence of A or B with at least one amino acid substitution at a proteolytic cleavage site; and
- F) a polypeptide comprising the amino acid sequence of A or B with at least one amino acid substitution at a cysteine residue.

A DNA according to claim 96, wherein said polypeptide includes at least one additional amino acid at the amino-terminus, at the carboxyl-terminus, or at both the amino-terminus and at the carboxyl-terminus.

A DNA according to claim 97, wherein said polypeptide includes at least one additional amino acid at the amino-terminus and at the carboxyl-terminus.

A DNA according to claim 97, wherein said polypeptide includes at least one additional amino acid at the amino-terminus.

A DNA according to claim 99, wherein said polypeptide includes a methionine at the amino-terminus.

A DNA according to claim 100, wherein said polypeptide includes at least one additional amino acid at the carboxyl-terminus.

A nucleic acid that hybridizes to a DNA complementary to the DNA defined in claim 96 under conditions of low stringency such that the nucleic acid is useful as a hybridization probe to detect DNA encoding the polypeptide of A or B.

of claims 15, 11, 14, 75, 15 or 95,

A vector comprising a DNA sequence coding for a TNF binding protein which binds TNF.

J. A vector comprising a DNA sequence defined in claim 96.

23. A vector according to claim 104, which is replicable in a prokaryotic and/or a eukaryotic host cell.

33. A vector according to claim 103, which is replicable in a prokaryotic cell.

With the sequence according to claim 106, wherein said DNA sequence encodes ATG at the amino terminus of the peptide.

136 18. A vector according to claim 186, which is replicable in Escherichia coli.

A vector according to claim 135, which is replicable in a eukaryotic cell.

37 10. A vector according to claim 109, which is replicable in a mammalian cell.

A vector according to claim 110, which is replicable in a Chinese Hamster Ovary

cell.

A vector according to claim 110, which is replicable in a COS cell.

A host cell containing a recombinant DNA molecule comprising a DNA sequence

defined in claim 97.

41. A host cell according to claim 113, which is a prokaryotic cell.

1/5. A host cell according to claim 1/4, which is Escherichia coli.

115. A host cell according to claim 113, which is a eukaryotic cell.

A host cell according to claim 116, which is a mammalian cell.

A host cell according to claim 117, which is a Chinese Hamster Ovary cell.

A,host cell according to claim, which is a COS cell.

A recombinant host cell according to claim 76, wherein the DNA molecule comprises promoter DNA, other than the promoter DNA for the native polypeptide having the ability to bind TNF, operatively linked to DNA sequence defined in claim 96 or 131

A process for preparing a polypeptide having the ability to bind TNF comprising producing the polypeptide in a recombinant host cell according to claim 120 under suitable conditions to express the DNA molecule contained therein to produce the polypeptide, and recovering the polypeptide.

A process for preparing a polypeptide having the ability to bind TNF comprising producing the polypeptide in a recombinant host cell according to claim under suitable conditions to express the DNA molecule contained therein to produce the polypeptide, and recovering the polypeptide.

A process according to claim 122, further comprising combining the recovered recombinant polypeptide with a pharmaceutically acceptable carrier to form a pharmaceutical composition.

A process according to claim 133, further comprising chemically derivatizing the harvested recombinant polypeptide.

A process according to claim 124, further comprising combining the chemically derivatized recombinant polypeptide with a pharmaceutically acceptable carrier to form a pharmaceutical composition.

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the isolated

The isolated

wherein said-recombinant DNA molecule is contained in an expression vector.

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An isolated DNA molecule according to one of claims 65, 71, 74, 75, 87, 95 or 36 wherein said polypeptide includes at least one additional amino acid at the amino-terminus, at the carboxyl-terminus, or at both the amino-terminus and at the carboxyl-terminus.

An isolated DNA molecule according to claim 127, wherein said polypeptide includes at least one additional amino acid at the amino-terminus and at the carboxyl-terminus.

An isolated DNA molecule according to claim 121, wherein said polypeptide includes at least one additional amino acid at the amino-terminus.

An isolated DNA molecule according to claim 129, wherein said polypeptide includes a methionine at the amino-terminus.

includes at least one additional amino acid at the carboxyl-terminus.

An isolated DNA molecule according to claim 65, 71, 74, 75, 81, 95 or 96, wherein said polypeptide includes a methionine at the amino-terminus and said amino acid substitution is at a glycosylation site.

An isolated DNA molecule according to claim 68, 71, 74, 75, 87, 93 or 96, wherein said polypeptide includes an amino acid substitution is at a glycosylation site.